1/16

Organism	gi number	Gene size & distance	Dis.(bp)
A. actinomycetemcomitans			-1
A. tumefaciens	17740473; 17740472		-4
B. capecia	•		58
C. burnetii			5
E. coli	1788328; 3915522		-1
E. faeciumpRUM plasmid	28849326; 28849325		-5
F. tularensis pFNL10 plasmid	4325240		-1
K. pneumoniae			57
M. bovis			65
M. tuberculosis	15610493; 7477590		-4
N. europaea A	; 22954686		-1
N. europaea B	22955251		10
N. europaea C			47
Nostos sp.PCC 7120	17135150; 17135149		-11
P. flourescens	; 23062981	F ,54	2
P. putida	24984546; 24984545		21
P. syringae	28850803; 28850802		29
R. conorii	15892044		80
S. typhimurium	16422119; 16422118		-1
S. aureus	14248182; 15925397	·	-1
S. pneumoniae	14973237; 14973236		3
A. coelicolor	4469319; 4469318		-1
S. viridochromogenes	15077435; 15077436		-4
Synechocystis sp. PCC 7942	22002543; 22002544		-5
Synechocystis sp. PCC 6803 A	16331058;		-4
Synechocystis sp. PCC 6803 B	16330633;		-1
T. ferrooxidans			80
Y. enterocolitica A			5
Y. enterocolitica B			56
		200	-

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Fig. 1

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A. actinomycetemcomitans

F. tularensis

S. aureus

S. pneumoniae

E. faecium

R. conorii

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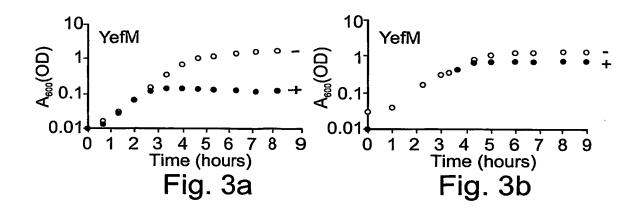
Synechocystis sp. PCC 7942

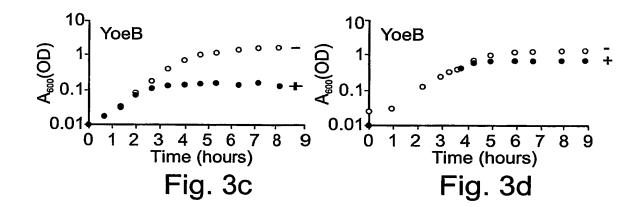
N. europaea B

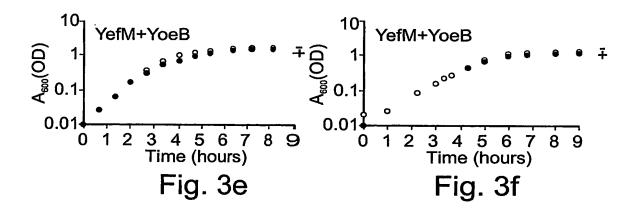
M. fuberculosis

C. burnetii

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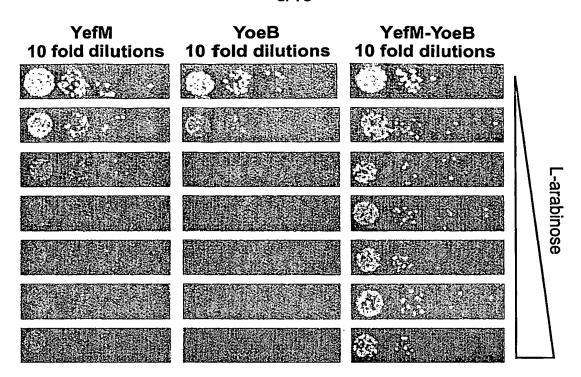
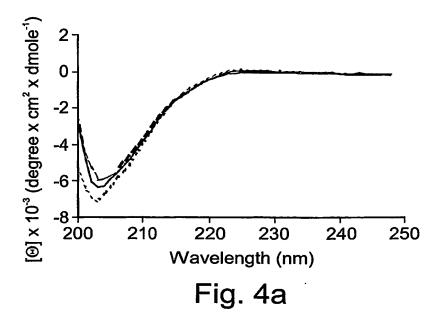
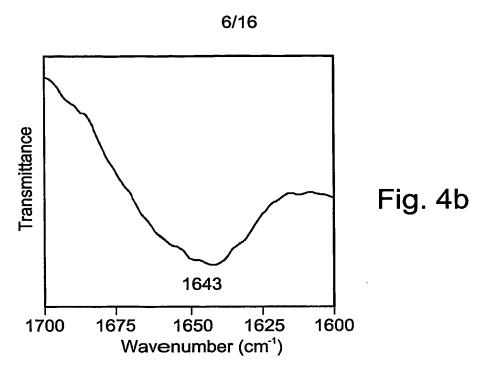
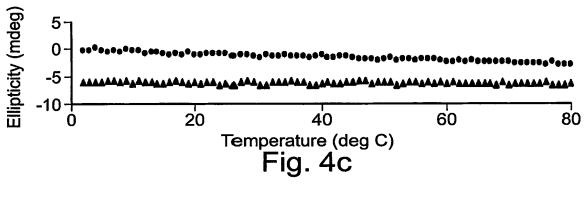


Fig. 3g







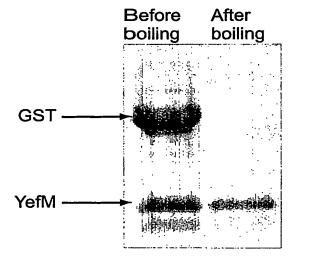
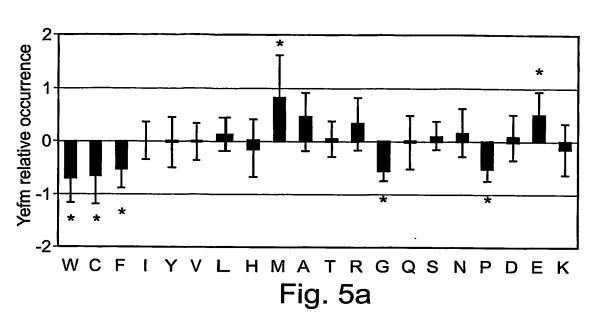


Fig. 4d





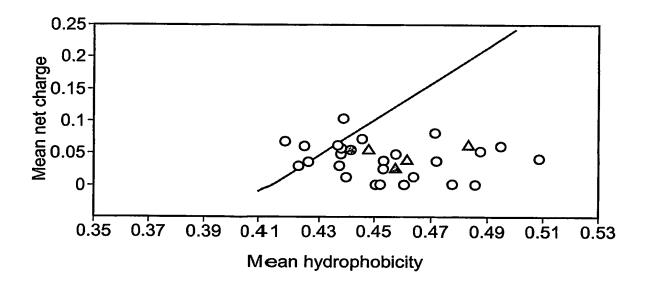
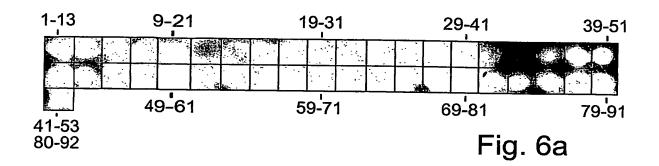
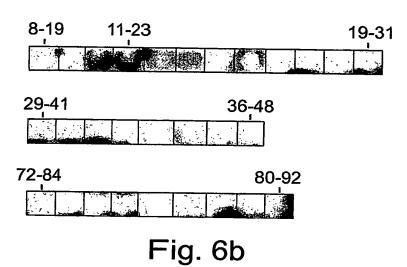


Fig. 5b

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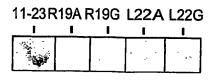


Fig. 6c

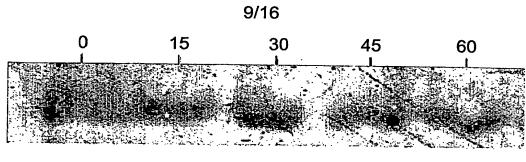
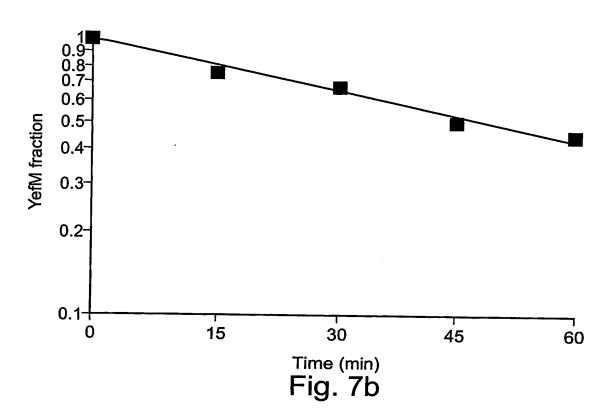
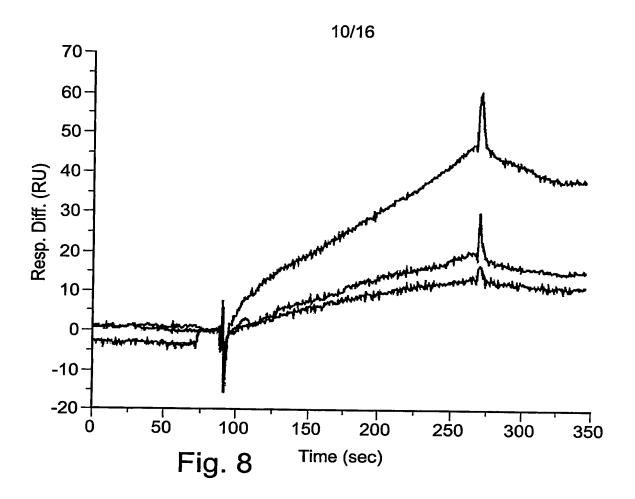


Fig. 7a





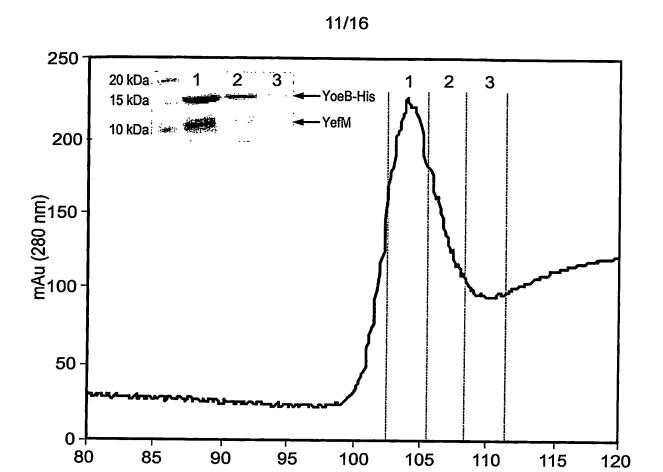


Fig. 9a

RT (min)



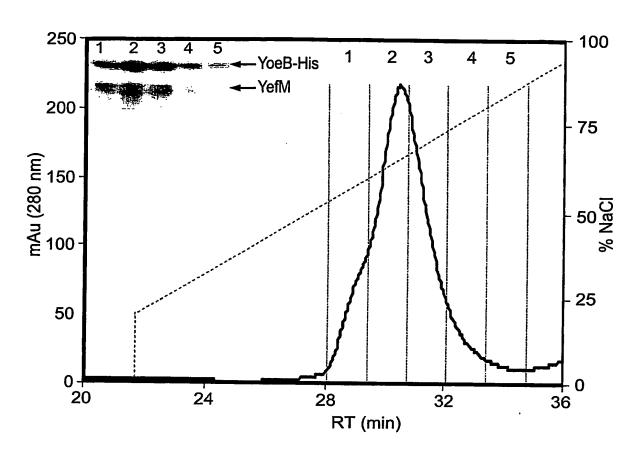


Fig. 9b



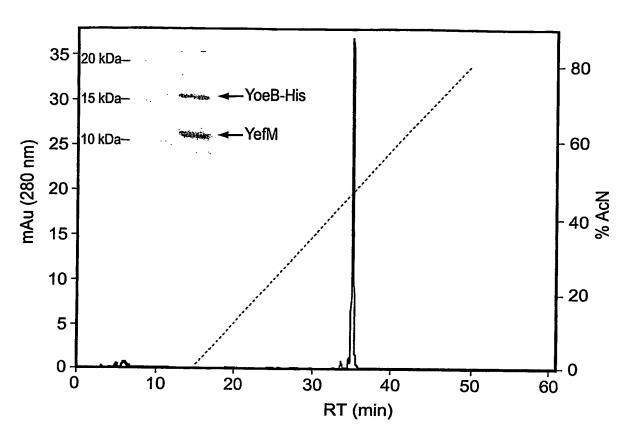
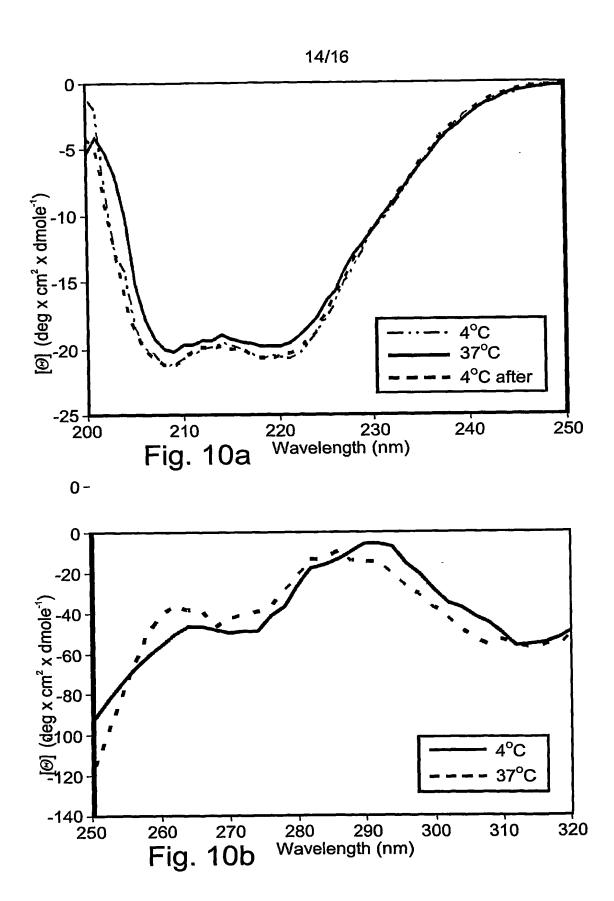
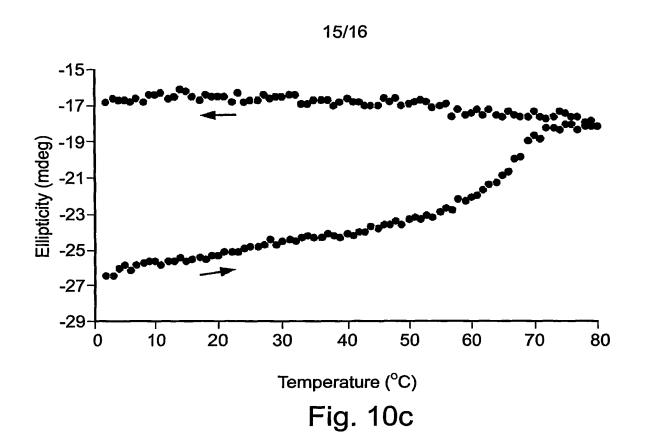
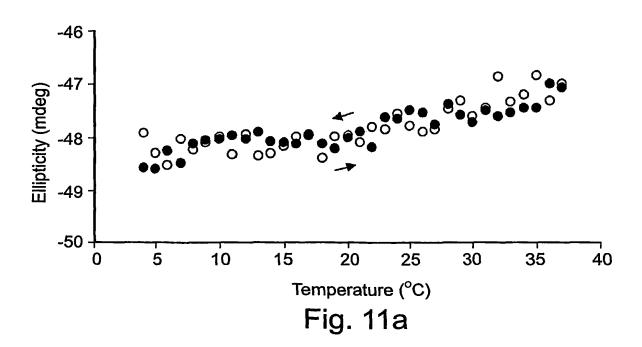


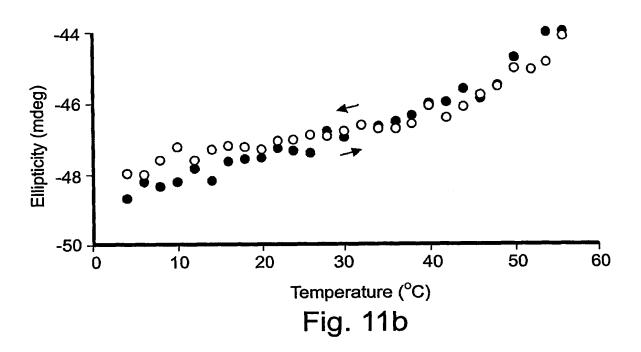
Fig. 9c

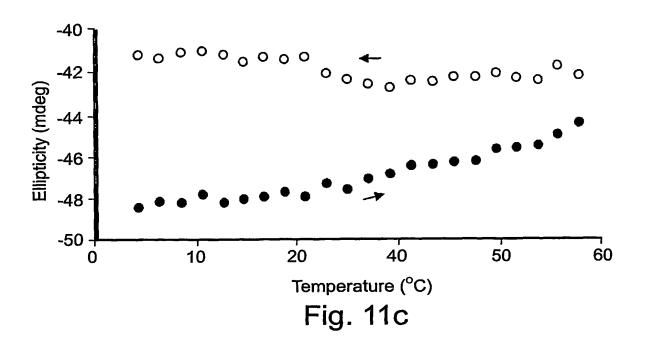






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